

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 09/889,330
Source: 1FW16
Date Processed by STIC: 1/14/05

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 01/14/2005

PATENT APPLICATION: US/09/889,330

TIME: 13:00:17

Input Set : A:\249146us.app

Output Set: N:\CRF4\01142005\I889330.raw

3 <110> APPLICANT: KOLTERMAN, ORVILLE G.
 4 YOUNG, ANDREW A.
 6 <120> TITLE OF INVENTION: NOVEL EXENDIN AGONIST FORMULATIONS AND METHODS OF
 ADMINISTRATION

7 THEREOF

9 <130> FILE REFERENCE: 249/146US

11 <140> CURRENT APPLICATION NUMBER: 09/889,330

12 <141> CURRENT FILING DATE: 2001-12-27

14 <150> PRIOR APPLICATION NUMBER: PCT/US00/00902

15 <151> PRIOR FILING DATE: 2000-01-14

17 <150> PRIOR APPLICATION NUMBER: US 60/116,380

18 <151> PRIOR FILING DATE: 1999-01-14

20 <150> PRIOR APPLICATION NUMBER: US 60/175,365

21 <151> PRIOR FILING DATE: 2000-01-10

23 <160> NUMBER OF SEQ ID NOS: 189

25 <170> SOFTWARE: PatentIn Ver. 3.2

27 <210> SEQ ID NO: 1

28 <211> LENGTH: 39

29 <212> TYPE: PRT

30 <213> ORGANISM: Heloderma horridum

32 <220> FEATURE:

33 <221> NAME/KEY: MOD_RES

34 <222> LOCATION: (39)

35 <223> OTHER INFORMATION: AMIDATION, Position 39 is Ser-NH2

37 <400> SEQUENCE: 1

38 His Ser Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu

39 1 5 10 15

41 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser

42 20 25 30

44 Ser Gly Ala Pro Pro Pro Ser

45 35

48 <210> SEQ ID NO: 2

49 <211> LENGTH: 39

50 <212> TYPE: PRT

51 <213> ORGANISM: Heloderma suspectum

53 <220> FEATURE:

54 <221> NAME/KEY: MOD_RES

55 <222> LOCATION: (39)

56 <223> OTHER INFORMATION: AMIDATION, Position 39 is Ser-NH2

58 <400> SEQUENCE: 2

59 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu

60 1 5 10 15

62 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser

63 20 25 30

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65 Ser Gly Ala Pro Pro Pro Ser
66 35
69 <210> SEQ ID NO: 3
70 <211> LENGTH: 39
71 <212> TYPE: PRT
72 <213> ORGANISM: Artificial Sequence
74 <220> FEATURE:
75 <223> OTHER INFORMATION: synthetic construct
77 <220> FEATURE:
78 <221> NAME/KEY: VARIANT
79 <222> LOCATION: (1)
80 <223> OTHER INFORMATION: His, Arg or Thr
82 <220> FEATURE:
83 <221> NAME/KEY: VARIANT
84 <222> LOCATION: (2)
85 <223> OTHER INFORMATION: Ser, Gly, Ala or Thr
87 <220> FEATURE:
88 <221> NAME/KEY: VARIANT
89 <222> LOCATION: (3)
90 <223> OTHER INFORMATION: Asp or Glu
92 <220> FEATURE:
93 <221> NAME/KEY: VARIANT
94 <222> LOCATION: (6)
95 <223> OTHER INFORMATION: Phe, Tyr or naphthalanine
97 <220> FEATURE:
98 <221> NAME/KEY: VARIANT
99 <222> LOCATION: (7)
100 <223> OTHER INFORMATION: Thr or Ser
102 <220> FEATURE:
103 <221> NAME/KEY: VARIANT
104 <222> LOCATION: (8)
105 <223> OTHER INFORMATION: Ser or Thr
107 <220> FEATURE:
108 <221> NAME/KEY: VARIANT
109 <222> LOCATION: (9)
110 <223> OTHER INFORMATION: Asp or Glu
112 <220> FEATURE:
113 <221> NAME/KEY: VARIANT
114 <222> LOCATION: (10)
115 <223> OTHER INFORMATION: Leu, Ile, Val, pentylglycine or Met
117 <220> FEATURE:
118 <221> NAME/KEY: VARIANT
119 <222> LOCATION: (14)
120 <223> OTHER INFORMATION: Leu, Ile, pentylglycine, Val or Met
122 <220> FEATURE:
123 <221> NAME/KEY: VARIANT
124 <222> LOCATION: (22)
125 <223> OTHER INFORMATION: Phe, Tyr or naphthalanine
127 <220> FEATURE:

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128 <221> NAME/KEY: VARIANT
129 <222> LOCATION: (23)
130 <223> OTHER INFORMATION: Ile, Val, Leu, pentylglycine, tert-butylglycine or Met
132 <220> FEATURE:
133 <221> NAME/KEY: VARIANT
134 <222> LOCATION: (24)
135 <223> OTHER INFORMATION: Glu or Asp
137 <220> FEATURE:
138 <221> NAME/KEY: VARIANT
139 <222> LOCATION: (25)
140 <223> OTHER INFORMATION: Trp, Phe, Tyr or naphthylalanine
142 <220> FEATURE:
143 <221> NAME/KEY: VARIANT
144 <222> LOCATION: (31)
145 <223> OTHER INFORMATION: independently Pro, homoproline, 3-hydroxyproline,
146     4-hydroxyproline, thioproline, N-alkylglycine,
147     N-alkylpentylglycine or N-alkylalanine
149 <220> FEATURE:
150 <221> NAME/KEY: VARIANT
151 <222> LOCATION: (36)..(38)
152 <223> OTHER INFORMATION: independently Pro, homoproline, 3-hydroxyproline,
153     4-hydroxyproline, thioproline, N-alkylglycine,
154     N-alkylpentylglycine or N-alkylalanine
156 <220> FEATURE:
157 <221> NAME/KEY: VARIANT
158 <222> LOCATION: (39)
159 <223> OTHER INFORMATION: Ser, Thr or Tyr
161 <220> FEATURE:
162 <223> OTHER INFORMATION: c-term is -OH or NH2, with the proviso that the compound
163     does not have the formula of either SEQ ID NOS 1 or 2
165 <400> SEQUENCE: 3
W--> 166 Xaa Xaa Xaa Gly Thr Xaa Xaa Xaa Xaa Xaa Ser Lys Gln Xaa Glu Glu
      167 1           5           10           15
      169 Glu Ala Val Arg Leu Xaa Xaa Xaa Xaa Leu Lys Asn Gly Gly Xaa Ser
      170           20           25           30
      172 Ser Gly Ala Xaa Xaa Xaa Xaa
      173           35
      175 <210> SEQ ID NO: 4
      176 <211> LENGTH: 38
      177 <212> TYPE: PRT
      178 <213> ORGANISM: Artificial Sequence
      180 <220> FEATURE:
      181 <223> OTHER INFORMATION: synthetic construct
      183 <220> FEATURE:
      184 <221> NAME/KEY: VARIANT
      185 <222> LOCATION: (1)
      186 <223> OTHER INFORMATION: His, Arg, or Tyr
      188 <220> FEATURE:
      189 <221> NAME/KEY: VARIANT

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Input Set : A:\249146us.app

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190 <222> LOCATION: (2)
191 <223> OTHER INFORMATION: Ser, Gly, Ala or Thr
193 <220> FEATURE:
194 <221> NAME/KEY: VARIANT
195 <222> LOCATION: (3)
196 <223> OTHER INFORMATION: Asp or Glu
198 <220> FEATURE:
199 <221> NAME/KEY: VARIANT
200 <222> LOCATION: (5)
201 <223> OTHER INFORMATION: Ala or Thr
203 <220> FEATURE:
204 <221> NAME/KEY: VARIANT
205 <222> LOCATION: (6)
206 <223> OTHER INFORMATION: Ala, Phe, Tyr or naphthylalanine
208 <220> FEATURE:
209 <221> NAME/KEY: VARIANT
210 <222> LOCATION: (7)
211 <223> OTHER INFORMATION: Thr or Ser
213 <220> FEATURE:
214 <221> NAME/KEY: VARIANT
215 <222> LOCATION: (8)
216 <223> OTHER INFORMATION: Ala, Ser or Thr
218 <220> FEATURE:
219 <221> NAME/KEY: VARIANT
220 <222> LOCATION: (9)
221 <223> OTHER INFORMATION: Asp or Glu
223 <220> FEATURE:
224 <221> NAME/KEY: VARIANT
225 <222> LOCATION: (10)
226 <223> OTHER INFORMATION: Ala, Leu, Ile, Val, pentylglycine or Met
228 <220> FEATURE:
229 <221> NAME/KEY: VARIANT
230 <222> LOCATION: (11)
231 <223> OTHER INFORMATION: Ala or Ser
233 <220> FEATURE:
234 <221> NAME/KEY: VARIANT
235 <222> LOCATION: (12)
236 <223> OTHER INFORMATION: Ala or Lys
238 <220> FEATURE:
239 <221> NAME/KEY: VARIANT
240 <222> LOCATION: (13)
241 <223> OTHER INFORMATION: Ala or Gln
243 <220> FEATURE:
244 <221> NAME/KEY: VARIANT
245 <222> LOCATION: (14)
246 <223> OTHER INFORMATION: Ala, Leu, Ile, pentylglycine, Val or Met
248 <220> FEATURE:
249 <221> NAME/KEY: VARIANT
250 <222> LOCATION: (15) .. (17)

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Input Set : A:\249146us.app

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251 <223> OTHER INFORMATION: Ala or Glu
253 <220> FEATURE:
254 <221> NAME/KEY: VARIANT
255 <222> LOCATION: (19)
256 <223> OTHER INFORMATION: Ala or Val
258 <220> FEATURE:
259 <221> NAME/KEY: VARIANT
260 <222> LOCATION: (20)
261 <223> OTHER INFORMATION: Ala or Arg
263 <220> FEATURE:
264 <221> NAME/KEY: VARIANT
265 <222> LOCATION: (21)
266 <223> OTHER INFORMATION: Ala or Leu
268 <220> FEATURE:
269 <221> NAME/KEY: VARIANT
270 <222> LOCATION: (22)
271 <223> OTHER INFORMATION: Ala, Phe, Tyr or naphthylalanine
273 <220> FEATURE:
274 <221> NAME/KEY: VARIANT
275 <222> LOCATION: (23)
276 <223> OTHER INFORMATION: Ile, Val, Leu, pentylglycine, tert-butylglycine or Met
278 <220> FEATURE:
279 <221> NAME/KEY: VARIANT
280 <222> LOCATION: (24)
281 <223> OTHER INFORMATION: Ala, Glu or Asp
283 <220> FEATURE:
284 <221> NAME/KEY: VARIANT
285 <222> LOCATION: (25)
286 <223> OTHER INFORMATION: Ala, Trp, Phe, Tyr or naphthylalanine
288 <220> FEATURE:
289 <221> NAME/KEY: VARIANT
290 <222> LOCATION: (26)
291 <223> OTHER INFORMATION: Ala or Leu
293 <220> FEATURE:
294 <221> NAME/KEY: VARIANT
295 <222> LOCATION: (27)
296 <223> OTHER INFORMATION: Ala or Lys
298 <220> FEATURE:
299 <221> NAME/KEY: VARIANT
300 <222> LOCATION: (28)
301 <223> OTHER INFORMATION: Ala or Asn
303 <220> FEATURE:
304 <221> NAME/KEY: MOD_RES
305 <222> LOCATION: (31)
306 <223> OTHER INFORMATION: Pro, homoproline, 3Hyp, 4Hyp, thioproline, N-alkylglycine,
307 N-alkylpentylglycine or N-alkylalanine
309 <220> FEATURE:
310 <221> NAME/KEY: MOD_RES
311 <222> LOCATION: (36)..(38)

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/889,330

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Input Set : A:\249146us.app
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; Xaa Pos. ~~1,2,3,5,7,8,9,10,14,22,23,24,25,31,36,37,38,39~~
Seq#:4; Xaa Pos. ~~1,2,3,5,6,7,8,9,10,11,12,13,14,15,16,17,19,20,21,22,23,24~~
Seq#:4; Xaa Pos. ~~25,26,27,28,31,36,37,38~~
Seq#:5; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,19,20,21,22,23
Seq#:5; Xaa Pos. 24,25,26,27,28,31,36,37,38,39
Seq#:15; Xaa Pos. 6
Seq#:20; Xaa Pos. 10
Seq#:21; Xaa Pos. 10
Seq#:22; Xaa Pos. 14
Seq#:23; Xaa Pos. 14
Seq#:24; Xaa Pos. 22
Seq#:27; Xaa Pos. 23
Seq#:28; Xaa Pos. 23
Seq#:31; Xaa Pos. 31,36,37,38
Seq#:32; Xaa Pos. 36,37,38
Seq#:33; Xaa Pos. 31,36,37,38
Seq#:34; Xaa Pos. 36,37,38
Seq#:35; Xaa Pos. 31,36,37,38
Seq#:36; Xaa Pos. 31,36,37,38
Seq#:37; Xaa Pos. 31,36,37,38
Seq#:38; Xaa Pos. 36,37,38
Seq#:39; Xaa Pos. 31,36,37,38
Seq#:81; Xaa Pos. 31,36,37,38
Seq#:82; Xaa Pos. 36,37,38
Seq#:83; Xaa Pos. 31
Seq#:84; Xaa Pos. 31,36,37
Seq#:85; Xaa Pos. 31,36,37
Seq#:86; Xaa Pos. 31,36
Seq#:89; Xaa Pos. 6
Seq#:93; Xaa Pos. 10
Seq#:94; Xaa Pos. 22
Seq#:95; Xaa Pos. 23
Seq#:99; Xaa Pos. 31,36,37
Seq#:115; Xaa Pos. 6
Seq#:116; Xaa Pos. 6
Seq#:127; Xaa Pos. 10
Seq#:128; Xaa Pos. 10
Seq#:137; Xaa Pos. 14
Seq#:138; Xaa Pos. 14
Seq#:151; Xaa Pos. 22
Seq#:152; Xaa Pos. 22
Seq#:155; Xaa Pos. 23
Seq#:156; Xaa Pos. 23
Seq#:181; Xaa Pos. 31,36,37,38

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Input Set : A:\249146us.app

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Seq#:182; Xaa Pos. 36,37,38

Seq#:183; Xaa Pos. 31,36,37

Seq#:184; Xaa Pos. 31,36

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:5; Line(s) 484

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/889,330

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Input Set : A:\249146us.app

Output Set: N:\CRF4\01142005\I889330.raw

L:166 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
M:341 Repeated in SeqNo=3
L:320 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
M:341 Repeated in SeqNo=4
L:481 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
M:341 Repeated in SeqNo=5
L:711 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:836 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:865 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:894 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:923 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:955 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:16
L:1032 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:16
L:1061 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:16
L:1143 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:16
M:341 Repeated in SeqNo=31
L:1175 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:32
L:1206 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:16
M:341 Repeated in SeqNo=33
L:1238 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:32
L:1269 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:16
M:341 Repeated in SeqNo=35
L:1303 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:16
M:341 Repeated in SeqNo=36
L:1337 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:16
M:341 Repeated in SeqNo=37
L:1369 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:32
L:1400 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:16
M:341 Repeated in SeqNo=39
L:2331 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81 after pos.:16
M:341 Repeated in SeqNo=81
L:2363 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82 after pos.:32
L:2389 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83 after pos.:16
L:2423 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84 after pos.:16
M:341 Repeated in SeqNo=84
L:2457 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:85 after pos.:16
M:341 Repeated in SeqNo=85
L:2491 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86 after pos.:16
M:341 Repeated in SeqNo=86
L:2562 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:89 after pos.:0
L:2651 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:93 after pos.:0
L:2680 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94 after pos.:16
L:2706 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:95 after pos.:16
L:2803 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:99 after pos.:16
M:341 Repeated in SeqNo=99
L:3144 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:115 after pos.:0
L:3170 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:116 after pos.:0
L:3406 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:127 after pos.:0

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L:3432 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:128 after pos.:0

L:3626 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:137 after pos.:0